



#12

SEQUENCE LISTING

<110> KNAUF, VIC C.
KRIDL, JEAN C.

<120> METHODS AND COMPOSITIONS FOR REGULATED TRANSCRIPTION
AND EXPRESSION OF HETEROLOGOUS GENES

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<141> 2001-02-12

<150> US 09/232,861

<151> 1999-01-15

<150> US 08/812,665

<151> 1997-03-07

<150> US 08/484,941

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<150> US 08/105,852

<151> 1993-08-10

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<150> US 07/168,190

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<150> US 07/054,369

<151> 1987-05-26

<150> US 07/742,834

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ctc	att	tct	ttg	tct	gaa	caa	gaq	ctt	qtc	qac	tqc	qac	aca	aac	gat	580

Leu	Ile	Ser	Leu	Ser	Glu	Gln	Glu	Leu	Val	Asp	Cys	Asp	Thr	Asn	Asp		
			175					180					185				
gat	ggc	tgc	atg	ggc	ggg	tac	atg	aat	agt	gcg	ttt	aac	tac	aca	atg	628	
Asp	Gly	Cys	Met	Gly	Gly	Tyr	Met	Asn	Ser	Ala	Phe	Asn	Tyr	Thr	Met		
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act	act	ggc	ggc	tta	acc	tct	gaa	tca	aat	tat	cct	tat	aaa	agc	aca	676	
Thr	Thr	Gly	Gly	Leu	Thr	Ser	Glu	Ser	Asn	Tyr	Pro	Tyr	Lys	Ser	Thr		
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Asp	Gly	Thr	Cys	Asn	Phe	Asn	Lys	Thr	Lys	Gln	Ile	Ala	Thr	Ser	Ile		
220					225					230					235		
aaa	ggg	ttt	gag	gat	gtc	ccg	gct	aac	gat	gag	aaa	gcc	cta	atg	aag	772	
Lys	Gly	Phe	Glu	Asp	Val	Pro	Ala	Asn	Asp	Glu	Lys	Ala	Leu	Met	Lys		
				240					245					250			
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Ala	Val	Ala	His	His	Pro	Val	Ser	Ile	Gly	Ile	Ala	Gly	Gly	Asp	Ile		
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Gly	Phe	Gln	Phe	Tyr	Ser	Ser	Gly	Val	Phe	Ser	Gly	Glu	Cys	Thr	Thr		
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His	Leu	Asp	His	Gly	Val	Thr	Ala	Val	Gly	Tyr	Gly	Arg	Ser	Lys	Asn		
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<211> 344

<212> PRT

<213> Brassica campestris

<400> 7

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Ala Met Gln Lys Arg His Ala Glu Trp Met Thr Glu His Gly Arg Val	35	40	45
Tyr Ala Asp Ala Asn Glu Lys Asn Asn Arg Tyr Ala Val Phe Lys Arg	50	55	60
Asn Val Glu Arg Ile Glu Arg Leu Asn Asp Val Gln Ser Gly Leu Thr	65	70	75
Phe Lys Leu Ala Val Asn Gln Phe Ala Asp Leu Thr Asn Glu Glu Phe	85	90	95
Arg Ser Met Tyr Thr Gly Phe Lys Gly Asn Ser Val Leu Ser Ser Arg	100	105	110
Thr Lys Pro Thr Ser Phe Arg Tyr Gln Asn Val Ser Ser Asp Ala Leu	115	120	125
Pro Val Ser Val Asp Trp Arg Lys Lys Gly Ala Val Thr Pro Ile Lys	130	135	140
Asp Gln Gly Leu Cys Gly Ser Cys Trp Ala Phe Ser Ala Val Ala Ala	145	150	155
Ile Glu Gly Val Ala Gln Ile Lys Lys Gly Lys Leu Ile Ser Leu Ser	165	170	175
Glu Gln Glu Leu Val Asp Cys Asp Thr Asn Asp Asp Gly Cys Met Gly	180	185	190
Gly Tyr Met Asn Ser Ala Phe Asn Tyr Thr Met Thr Thr Gly Gly Leu	195	200	205
Thr Ser Glu Ser Asn Tyr Pro Tyr Lys Ser Thr Asp Gly Thr Cys Asn	210	215	220
Phe Asn Lys Thr Lys Gln Ile Ala Thr Ser Ile Lys Gly Phe Glu Asp	225	230	235
Val Pro Ala Asn Asp Glu Lys Ala Leu Met Lys Ala Val Ala His His	245	250	255
Pro Val Ser Ile Gly Ile Ala Gly Gly Asp Ile Gly Phe Gln Phe Tyr	260	265	270
Ser Ser Gly Val Phe Ser Gly Glu Cys Thr Thr His Leu Asp His Gly	275	280	285
Val Thr Ala Val Gly Tyr Gly Arg Ser Lys Asn Gly Leu Lys Tyr Trp	290	295	300
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<213> Lycopersicon esculentum

<400> 10

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              20              25              30
Val Met Ala Leu Arg Asp Ile Pro Pro Gln Glu Thr Leu Leu Lys Met
      35              40              45
Lys Leu Leu Pro Thr Asn Ile Leu Gly Leu Cys Asn Glu Pro Cys Ser
      50              55              60
Ser Asn Ser Asp Cys Ile Gly Ile Thr Leu Cys Gln Phe Cys Lys Glu
      65              70              75              80
Lys Thr Asp Gln Tyr Gly Leu Thr Tyr Arg Thr Cys Asn Leu Leu Pro
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<210> 11

<211> 14

<212> PRT

<213> Lycopersicon esculentum

<400> 11

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<210> 12

<211> 18

<212> PRT

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Protease
inhibitor PA1b peptide sequence

<400> 12

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 1              5              10              15
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Cys Ile

<210> 13

<211> 13

<212> PRT

<213> Pisum sp.

<400> 13

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<210> 14
<211> 13
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<213> Phaseolus limensis

<400> 14
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1 5 10

<210> 15
<211> 12
<212> PRT
<213> Homo sapiens

<400> 15
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1 5 10

<210> 16
<211> 18
<212> PRT
<213> Lycopersicon esculentum

<400> 16
Thr Asn Ile Leu Gly Leu Cys Asn Glu Pro Cys Ser Ser Asn Ser Asp
1 5 10 15

Cys Ile

<210> 17
<211> 18
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<213> Unknown Organism

<220>
<223> Description of Unknown Organism: Protease
inhibitor PA1b peptide sequence

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Cys Arg

<210> 18
<211> 17
<212> PRT
<213> Hordeum vulgare

<400> 18
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1 5 10 15

Ala

<210> 19
<211> 16
<212> PRT
<213> Triticum sp.

<400> 19
Val Ser Ala Leu Thr Gly Cys Arg Ala Met Val Lys Leu Gln Cys Val
1 5 10 15

<210> 20
<211> 16
<212> PRT
<213> Triticum sp.

<400> 20
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1 5 10 15

<210> 21
<211> 20
<212> PRT
<213> Panicum miliaceum

<400> 21
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1 5 10 15

Thr Ala Cys Gly
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<210> 22
<211> 18
<212> PRT
<213> Ricinus communis

<400> 22
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1 5 10 15

Gly Gln

<210> 23
<211> 18
<212> PRT
<213> Unknown Organism

<220>
<223> Description of Unknown Organism: Napin small
subunit peptide sequence

<400> 23

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Gln Ser

<210> 24

<211> 4656

<212> DNA

<213> Lycopersicon esculentum

<220>

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<222> (1379) .. (1444)

<220>

<221> CDS

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<211> 96

<212> PRT

<213> Lycopersicon esculentum

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			20					25					30		

Val	Met	Ala	Leu	Arg	Asp	Ile	Pro	Pro	Gln	Glu	Thr	Leu	Leu	Lys	Met
		35					40					45			

Lys	Leu	Leu	Pro	Thr	Asn	Ile	Leu	Gly	Leu	Cys	Asn	Glu	Pro	Cys	Ser
	50					55					60				

Ser	Asn	Ser	Asp	Cys	Ile	Gly	Ile	Thr	Leu	Cys	Gln	Phe	Cys	Lys	Glu
65					70					75					80

Lys	Thr	Asp	Gln	Tyr	Gly	Leu	Thr	Tyr	Arg	Thr	Cys	Asn	Leu	Leu	Pro
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<213> Lycopersicon esculentum

<400> 26

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30

<210> 28

<211> 48

<212> DNA

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<223> Description of Artificial Sequence: Synthetic
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48

<210> 29

<211> 16

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
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<210> 30

<211> 43

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
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<210> 31

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<212> DNA

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29

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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
peptide

<400> 33
Ala Ala Lys Pro Glu Thr Val Glu Lys Val
1 5 10

<210> 34
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 34
cttaagaagt aaccgggct gcagtttttag tattaagag

39

<210> 35
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide polylinker sequence

<400> 35
agctcgttac cgaattcgag ctcggtac

28